



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/696,699
Source: TFW
Date Processed by STIC: 11/7/03

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

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Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/efc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
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Revised 10/08/03



IFWO

RAW SEQUENCE LISTING

DATE: 11/07/2003

PATENT APPLICATION: US/10/696,699

TIME: 11:50:17

Input Set : A:\D6453CIPSEQ.txt

Output Set: N:\CRF4\11062003\J696699.raw

2 <110> APPLICANT: Sanders, Bob G.
 3 Kline, Kimberly
 4 Yu, Weiping
 5 Liu, Hui
 6 Hantash, Feras
 8 <120> TITLE OF INVENTION: Tocopherol Associated Protein and Uses Thereof
 10 <130> FILE REFERENCE: D6453CIP
 E--> 12 <140> CURRENT APPLICATION NUMBER: US/10/696,699
 12 <141> CURRENT FILING DATE: 2003-10-29
 13 <150> PRIOR APPLICATION NUMBER: US 10/419,629
 14 <151> PRIOR FILING DATE: 2003-04-21
 16 <160> NUMBER OF SEQ ID NOS: 19

ERRORED SEQUENCES

251 <210> SEQ ID NO: 8
 252 <211> LENGTH: 16
 253 <212> TYPE: PRT
 254 <213> ORGANISM: Homo sapiens
 W--> 255 <220> FEATURE:
 256 <221> NAME/KEY: PEPTIDE
 257 <223> OTHER INFORMATION: 16 amino acids from the c-terminus of TAP attached
 258 to keyhole limpet hemocyanin
 260 <400> SEQUENCE: 8
 E--> 261 Lys (Als) Ser Glu Glu Lys Met Lys Gln Leu Gly Ala Gly Thr Pro
 262 5 10 15
 263 Lys
 264 16

invalid amino acid designator
 delete '16' Per Sequence Rules, number the amino acids under every 5 amino acids

*Please correct this
 in all similarly numbered
 amino acid sequences*

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/696,699

DATE: 11/07/2003

TIME: 11:50:18

Input Set : A:\D6453CIPSEQ.txt

Output Set: N:\CRF4\11062003\J696699.raw

L:12 M:282 E: Numeric Field Identifier Missing, <140> CURRENT APPLICATION NUMBER: is Added.
L:25 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:1
L:57 M:283 W: Missing Blank Line separator, <220> field identifier
L:121 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:155 M:283 W: Missing Blank Line separator, <220> field identifier
L:219 M:283 W: Missing Blank Line separator, <220> field identifier
L:255 M:283 W: Missing Blank Line separator, <220> field identifier
L:261 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:1
L:335 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:14
L:362 M:283 W: Missing Blank Line separator, <220> field identifier
L:414 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:16
L:437 M:283 W: Missing Blank Line separator, <220> field identifier
L:481 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:18
L:500 M:283 W: Missing Blank Line separator, <220> field identifier